### AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions, and listings, of the claims in the application.

1. (Currently amended) A fusion polypeptide comprising a first polypeptide operably linked to a second polypeptide,

wherein the first polypeptide comprises a polypeptide sequence with at least 85% homology to an extracellular portion of a glycoprotein Ibα polypeptide of SEQ ID NO:1, provided said glycoprotein Ibα polypeptide includes an amino acid other than glycine at position 233 or other than methionine at position 239 relative to the amino acid sequence of a wild-type GPIbα polypeptide, and said first polypeptide binds a polypeptide selected from the group consisting of leukocyte integrin Mac-1 polypeptide, von Willebrand factor, thrombin and P-selectin; and

wherein the second polypeptide comprises at least a region of an immunoglobulin heavy chain polypeptide.

## 2. (Cancelled)

3. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds to at least two of the polypeptides selected from the group consisting of leukocyte integrin Mac-1 polypeptide, von Willebrand factor, thrombin and P-selectin.

#### 4. (Cancelled)

5. (Previously presented) The fusion polypeptide of claim 1, wherein said polypeptide comprises SEQ ID NO: 5.

- 6. (Previously presented) The fusion polypeptide of claim 1, wherein said fusion polypeptide is more resistant to proteolysis than a wild-type GP Ibα polypeptide.
- 7. (Original) The fusion polypeptide of claim 1, wherein said first polypeptide binds with higher affinity to a von Willebrand factor polypeptide than a wild-type glycoprotein Ibα polypeptide binds to said von Willebrand factor polypeptide.
- 8. (Original) The fusion polypeptide of claim 1, wherein said first polypeptide comprises at least one of the amino acid substitutions G233V or M239V relative to the amino acid sequence of a wild-type GPIbα polypeptide.
- 9. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide comprises the amino acid substitutions G233V and M239V relative to the amino acid sequence of a wild-type GPIbα polypeptide.

# 10. (Cancelled)

- 11. (Original) The fusion polypeptide of claim 1, wherein said second polypeptide comprises an Fc region of an immunoglobulin heavy chain.
- 12. (Original) The fusion polypeptide of claim 11, wherein said second polypeptide has less effector function than the effector function of a Fc region of a wild-type immunoglobulin heavy chain.
- 13. (Original) The fusion polypeptide of claim 12, wherein said second polypeptide binds with low or no affinity to a Fc receptor.
- 14. (Original) The fusion polypeptide of claim 12, wherein said second polypeptide binds with low or no affinity to complement protein C1q.

#### 15-19. (Cancelled)

- 20. (Currently amended) The fusion polypeptide of claim 1, wherein said fusion polypeptide comprises the amino acid sequence of GPIbα302/4X-Ig (SEQ ID NO:3), or GPIbα290/2V-Ig (SEQ ID NO:5).
- 21. (Original) A multimeric polypeptide comprising the fusion polypeptide of claim 1.
- 22. (Original) The multimeric polypeptide of claim 21, wherein said multimeric polypeptide is a dimer.

# 23-26. (Cancelled)

27. (Original) A pharmaceutical composition comprising the fusion polypeptide of claim 1.

## 28-53. (Cancelled)

- 54. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds at least three polypeptides selected from the group consisting of leukocyte integrin Mac-1 polypeptide, von Willebrand factor, thrombin and P-selectin.
- 55. (previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds leukocyte integrin Mac-1 polypeptide, von Willebrand factor, thrombin and P-selectin.
- 56. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds leukocyte integrin Mac-1 polypeptide.

## 57. (Cancelled)

- 58. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds thrombin.
- 59. (Previously presented) The fusion polypeptide of claim 1, wherein said first polypeptide binds P-selectin.
- 60. (Previously presented) A fusion polypeptide comprising a first polypeptide operably linked to a second polypeptide,

wherein the first polypeptide consists essentially of a polypeptide sequence with at least 85% homology to an extracellular portion of a glycoprotein Ibα polypeptide of SEQ ID NO:1, provided said glycoprotein Ibα polypeptide includes an amino acid other than glycine at position 233 or other than methionine at position 239 relative to the amino acid sequence of a wild-type GPIbα polypeptide and said first polypeptide binds von Willebrand factor polypeptide; and

wherein the second polypeptide consists essentially of an immunoglobulin heavy chain polypeptide, wherein said immunoglobulin heavy chain polypeptide comprises a Fc region.

## 61-62. (Cancelled)

- 63. (Previously presented) A polypeptide consisting essentially of the amino acid sequence of SEQ ID NO:1.
- 64. (Previously presented) The polypeptide of claim 63, wherein the amino acid sequence of the polypeptide consists of SEO ID NO:1.
- 65. (Previously presented) A polypeptide consisting essentially of the amino acid sequence of SEQ ID NO:5.

66. (Previously presented) The polypeptide of claim 65, wherein the amino acid sequence of the polypeptide consists of SEQ ID NO:5.